



## RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 101647,519  
Source: IFW0  
Date Processed by STIC: 2/13/04

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 703-308-4212; FAX: 703-308-4221

Effective 12/13/03: TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - cPAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry directly to (EFFECTIVE 12/01/03):  
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 10/08/03



IFWO

RAW SEQUENCE LISTING  
 PATENT APPLICATION: US/10/647,519

DATE: 02/13/2004  
 TIME: 12:04:44

Input Set : A:\2314-269seq.txt  
 Output Set: N:\CRF4\02132004\J647519.raw

3 <110> APPLICANT: Fainzilber, Michael M.  
 4 Kits, Karel S.  
 5 Burlingame, Alma L.  
 6 Olivera, Baldomero M.  
 7 Walker, Craig  
 8 Watkins, Maren  
 9 Shetty, Reshma  
 10 Cruz, Lourdes J.  
 11 Imperial, Julita  
 12 Colledge, Clark  
 13 University of Utah Research Foundation  
 14 Vrije Universiteit  
 15 Regents of the University of California  
 17 <120> TITLE OF INVENTION: Gamma-Conopeptides  
 19 <130> FILE REFERENCE: 2314-147-sq2  
 C--> 21 <140> CURRENT APPLICATION NUMBER: US/10/647,519  
 C--> 22 <141> CURRENT FILING DATE: 2003-08-26  
 24 <150> PRIOR APPLICATION NUMBER: US 60/069,706  
 25 <151> PRIOR FILING DATE: 1997-12-16  
 27 <160> NUMBER OF SEQ ID NOS: 47  
 29 <170> SOFTWARE: PatentIn Ver. 2.0  
 31 <210> SEQ ID NO: 1  
 32 <211> LENGTH: 42  
 33 <212> TYPE: PRT  
 34 <213> ORGANISM: Artificial Sequence  
 36 <220> FEATURE:  
 37 <223> OTHER INFORMATION: Description of Artificial Sequence:generic formula  
 38 of gamma-conopeptides  
 40 <220> FEATURE:  
 41 <221> NAME/KEY: PEPTIDE  
 42 <222> LOCATION: (1)..(13)  
 43 <223> OTHER INFORMATION: Xaa at residues 1, 2, 3, 4, 5, and 6 may be  
 44 des-Xaa or any amino acid; Xaa at residues 8, 9,  
 45 10, 11 and 12 may be any amino acid; Xaa at  
 46 residue 13 may be des-Xaa or any amino acid.  
 48 <220> FEATURE:  
 49 <221> NAME/KEY: PEPTIDE  
 50 <222> LOCATION: (15)..(19)  
 51 <223> OTHER INFORMATION: Xaa at residues 15, 16, 17 and 18 may be any amino  
 52 acid; Xaa at residue 19 is Glu,  
 53 gamma-carboxyglutamate or Gln.  
 55 <220> FEATURE:  
 56 <221> NAME/KEY: PEPTIDE

pg. 6  
 Does Not Comply  
 Corrected Diskette Needed  
 (pg. 7-9)



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TIME: 12:04:44

Input Set : A:\2314-269seq.txt

Output Set: N:\CRF4\02132004\J647519.raw

117 <222> LOCATION: (30)..(42)

118 <223> OTHER INFORMATION: Xaa at residues 30, 31 and 32 may be des-Xaa or

119 any amino acid; Xaa at residues 34, 35, 36, 37,

120 38, 39, 40, 41 and 42 may be des-Xaa or any amino

121 acid.

123 <400> SEQUENCE: 2

W--> 124 Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa

125 1 5 10 15

127 Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa

128 20 25 30

130 Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa

131 35 40

134 <210> SEQ ID NO: 3

135 <211> LENGTH: 39

136 <212> TYPE: PRT

137 <213> ORGANISM: Artificial Sequence

139 <220> FEATURE:

140 <223> OTHER INFORMATION: Description of Artificial Sequence:generic formula

141 of gamma-conopeptides

143 <220> FEATURE:

144 <221> NAME/KEY: PEPTIDE

145 <222> LOCATION: (1)..(13)

146 <223> OTHER INFORMATION: Xaa at residue 1 is any amino acid; Xaa at

147 residues 2, 3, 4, 5 and 6 may be des-Xaa or any

148 amino acid; Xaa at residues 8, 9, 10, 11, 12 and

149 13 may be any amino acid.

151 <220> FEATURE:

152 <221> NAME/KEY: PEPTIDE

153 <222> LOCATION: (27)..(39)

154 <223> OTHER INFORMATION: Xaa at residues 27, 28, 29, 31, 32, 33, 34, 35, 36

155 and 37 may be any amino acid; Xaa at residues 38

156 and 39 may be des-Xaa or any amino acid.

158 <220> FEATURE:

159 <221> NAME/KEY: PEPTIDE

160 <222> LOCATION: (15)..(19)

161 <223> OTHER INFORMATION: Xaa at residues 15, 16, 17 and 18 may be any amino

162 acid; Xaa at residue 19 is Glu or

163 gamma-carboxyglutamate.

165 <400> SEQUENCE: 3

W--> 166 Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa

167 1 5 10 15

169 Xaa Xaa Xaa Cys Cys Ser Asn Ser Cys Asp Xaa Xaa Xaa Cys Xaa Xaa

170 20 25 30

172 Xaa Xaa Xaa Xaa Xaa Xaa Xaa

173 35

176 <210> SEQ ID NO: 4

177 <211> LENGTH: 39

178 <212> TYPE: PRT

179 <213> ORGANISM: Artificial Sequence

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Input Set : A:\2314-269seq.txt

Output Set: N:\CRF4\02132004\J647519.raw

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181 <220> FEATURE:
182 <223> OTHER INFORMATION: Description of Artificial Sequence:generic
183     sequence of gamma-conopeptides.
185 <220> FEATURE:
186 <221> NAME/KEY: PEPTIDE
187 <222> LOCATION: (1)..(13)
188 <223> OTHER INFORMATION: Xaa at residue 1 is any amino acid; Xaa at
189     residues 2, 3, 4, 5 and 6 may be des-Xaa or any
190     amino acid; Xaa at residues 8, 9, 10, 11, 12 and
191     13 may be any amino acid.
193 <220> FEATURE:
194 <221> NAME/KEY: PEPTIDE
195 <222> LOCATION: (15)..(19)
196 <223> OTHER INFORMATION: Xaa at residue 15 is Ser or Thr; Xaa at residues
197     16, 17 and 18 may be any amino acid; Xaa at
198     residue 19 is Glu or gamma-carboxyglutamate.
200 <220> FEATURE:
201 <221> NAME/KEY: PEPTIDE
202 <222> LOCATION: (27)..(39)
203 <223> OTHER INFORMATION: Xaa at residues 27, 28, 29, 31, 32, 33, 34, 35, 36
204     and 37 may be any amino acid; Xaa at residues 38
205     and 39 may be des-Xaa or any amino acid.
207 <400> SEQUENCE: 4
W--> 208 Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa
      209   1           5           10           15
      211 Xaa Xaa Xaa Cys Cys Ser Asn Ser Cys Asp Xaa Xaa Xaa Cys Xaa Xaa
      212           20           25           30
      214 Xaa Xaa Xaa Xaa Xaa Xaa Xaa
      215           35
218 <210> SEQ ID NO: 5
219 <211> LENGTH: 34
220 <212> TYPE: PRT
221 <213> ORGANISM: Artificial Sequence
223 <220> FEATURE:
224 <223> OTHER INFORMATION: Description of Artificial Sequence:generic
225     sequence of gamma-conopeptides.
227 <220> FEATURE:
228 <221> NAME/KEY: PEPTIDE
229 <222> LOCATION: (1)..(6)
230 <223> OTHER INFORMATION: Xaa at residues 1 and 2 may be des-Xaa or any
231     amino acid; Xaa at residue 3 is Asp, Glu or
232     gamma-carboxyglutamate; Xaa at residues 5 and 6
233     may be any amino acid.
235 <220> FEATURE:
236 <221> NAME/KEY: PEPTIDE
237 <222> LOCATION: (7)..(16)
238 <223> OTHER INFORMATION: Xaa at residue 7 is Trp or 6-bromo-Trp; Xaa at
239     residues 9, 10, 13 and 14 may be any amino acid;
240     Xaa at residue 16 is Glu or

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Input Set : A:\2314-269seq.txt

Output Set: N:\CRF4\02132004\J647519.raw

241 gamma-carboxyglutamate.  
 243 <220> FEATURE:  
 244 <221> NAME/KEY: PEPTIDE  
 245 <222> LOCATION: (28)..(34)  
 246 <223> OTHER INFORMATION: Xaa at residues 28, 30, 31, 32, 33 and 34 may be  
 247 any amino acid.  
 249 <400> SEQUENCE: 5  
 W--> 250 Xaa Xaa Xaa Cys Xaa Xaa Xaa Phe Xaa Xaa Cys Thr Xaa Xaa Ser Xaa  
 251 1 5 10 15  
 253 Cys Cys Ser Asn Ser Cys Asp Gln Thr Tyr Cys Xaa Leu Xaa Xaa Xaa  
 254 20 25 30  
 256 Xaa Xaa  
 260 <210> SEQ ID NO: 6  
 261 <211> LENGTH: 32  
 262 <212> TYPE: PRT  
 263 <213> ORGANISM: Conus pennaceus  
 265 <220> FEATURE:  
 266 <221> NAME/KEY: PEPTIDE  
 267 <222> LOCATION: (1)..(31)  
 268 <223> OTHER INFORMATION: Xaa at residue 5 is Trp or 6-bromo-Trp; Xaa at  
 269 residues 14 and 26 are Glu or  
 270 gamma-carboxyglutamate; Xaa at residue 31 is Pro  
 271 or hydroxy-Pro.  
 273 <400> SEQUENCE: 6  
 W--> 274 Asp Cys Thr Ser Xaa Phe Gly Arg Cys Thr Val Asn Ser Xaa Cys Cys  
 275 1 5 10 15  
 277 Ser Asn Ser Cys Asp Gln Thr Tyr Cys Xaa Leu Tyr Ala Phe Xaa Ser  
 278 20 25 30  
 284 <210> SEQ ID NO: 7  
 285 <211> LENGTH: 34  
 286 <212> TYPE: PRT  
 287 <213> ORGANISM: Conus textile  
 289 <220> FEATURE:  
 290 <221> NAME/KEY: PEPTIDE  
 291 <222> LOCATION: (1)..(34)  
 292 <223> OTHER INFORMATION: Xaa at residues 1, 7 and 34 are Trp or  
 293 6-bromo-Trp; Xaa at residues 3 and 16 are Glu or  
 294 gamma-carboxyglutamate; Xaa at residues 31 and 32  
 295 are Pro or hydroxy-Pro.  
 297 <400> SEQUENCE: 7  
 W--> 298 Xaa Leu Xaa Cys Ser Val Xaa Phe Ser His Cys Thr Lys Asp Ser Xaa  
 299 1 5 10 15  
 301 Cys Cys Ser Asn Ser Cys Asp Gln Thr Tyr Cys Thr Leu Met Xaa Xaa  
 302 20 25 30  
 304 Asp Xaa  
 308 <210> SEQ ID NO: 8  
 309 <211> LENGTH: 39  
 310 <212> TYPE: PRT  
 311 <213> ORGANISM: Conus textile

RAW SEQUENCE LISTING ERROR SUMMARY  
 PATENT APPLICATION: US/10/647,519

DATE: 02/13/2004  
 TIME: 12:04:45

Input Set : A:\2314-269seq.txt  
 Output Set: N:\CRF4\02132004\J647519.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 1,2,3,4,5,6,8,9,10,11,12,13,15,16,17,18,19,22,23,24,25,27 ✓  
 Seq#:1; Xaa Pos. 28,29,30,31,32,34,35,36,37,38,39,40,41,42 ✓  
 Seq#:2; Xaa Pos. 1,2,3,4,5,8,9,10,11,12,13,15,16,17,18,19,22,23,24,25,27 ✓  
 Seq#:2; Xaa Pos. 28,29,30,31,32,34,35,36,37,38,39,40,41,42 ✓  
 Seq#:3; Xaa Pos. 1,2,3,4,5,6,8,9,10,11,12,13,15,16,17,18,19,27,28,29,31,32 ✓  
 Seq#:3; Xaa Pos. 33,34,35,36,37,38,39 ✓  
 Seq#:4; Xaa Pos. 1,2,3,4,5,8,9,10,11,12,13,15,16,17,18,19,27,28,29,31,32 ✓  
 Seq#:4; Xaa Pos. 33,34,35,36,37,38,39 ✓  
 Seq#:5; Xaa Pos. 1,2,3,5,6,7,9,10,13,14,16,28,30,31,32,33,34 ✓  
 Seq#:6; Xaa Pos. 5,14,26,34 ✓  
 Seq#:7; Xaa Pos. 1,3,7,16,31,32,34 ✓  
 Seq#:8; Xaa Pos. 1,2,4,10,19,31,34,36,37,39 ✓  
 Seq#:9; Xaa Pos. 9,13,17 ✓  
 Seq#:10; Xaa Pos. 2,3,10,12,18,26,32,33 ✓  
 Seq#:11; Xaa Pos. 3,5,16,18,22,25,31 ✓  
 Seq#:12; Xaa Pos. 1,5,8,11 ✓  
 Seq#:13; Xaa Pos. 5,8,10,15,16,23 ✓  
 Seq#:14; Xaa Pos. 4,9,12,13,17 ✓  
 Seq#:15; Xaa Pos. 8,15,16 ✓  
 Seq#:24; N Pos. 17 ✓  
 Seq#:36; Xaa Pos. 3,7,8,14,18 ✓  
 Seq#:37; Xaa Pos. 1 ✓  
 Seq#:39; N Pos. 12 ✓  
 Seq#:41; N Pos. 6,9,12,15,18,21,24 ✓  
 Seq#:42; Xaa Pos. 14,26,31 ✓  
 Seq#:43; Xaa Pos. 9,13 ✓  
 Seq#:44; Xaa Pos. 2 ✓  
 Seq#:45; Xaa Pos. 1,6,7,8,12,17,18,23 ✓

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<210> SEQ ID NO 24  
<211> LENGTH: 533  
<212> TYPE: DNA  
<213> ORGANISM: Conus textile  
<220> FEATURE:  
<221> NAME/KEY: CDS  
<222> LOCATION: (110)..(337)  
<400> SEQUENCE: 24

ctctgccggt tgacac<sup>o</sup>tca tctactctct cagtctccct gacagctgcc ttcagtcgac 60

*mandatory, please  
explain "N" locations.  
please see error  
explanation  
on page 8.*

The type of errors shown exist throughout  
the Sequence Listing. Please check subsequent  
sequences for similar errors.  
Seq. (39, 41)

## VARIABLE LOCATION SUMMARY

PATENT APPLICATION: US/10/647,519

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Input Set : A:\2314-269seq.txt

Output Set: N:\CRF4\02132004\J647519.raw

Use of n's or Xaa's (NEW RULES):

Use of n's and/or Xaa's have been detected in the Sequence Listing.

Use of <220> to <223> is MANDATORY if n's or Xaa's are present.

in &lt;220&gt; to &lt;223&gt; section, please explain location of n or Xaa, and which residue n or Xaa represents.

Error Explanation:

Seq#:1; Xaa Pos. 1,2,3,4,5,6,8,9,10,11,12,13,15,16,17,18,19,22,23,24,25,27  
 Seq#:1; Xaa Pos. 28,29,30,31,32,34,35,36,37,38,39,40,41,42  
 Seq#:2; Xaa Pos. 1,2,3,4,5,6,8,9,10,11,12,13,15,16,17,18,19,22,23,24,25,27  
 Seq#:2; Xaa Pos. 28,29,30,31,32,34,35,36,37,38,39,40,41,42  
 Seq#:3; Xaa Pos. 1,2,3,4,5,6,8,9,10,11,12,13,15,16,17,18,19,27,28,29,31,32  
 Seq#:3; Xaa Pos. 33,34,35,36,37,38,39  
 Seq#:4; Xaa Pos. 1,2,3,4,5,6,8,9,10,11,12,13,15,16,17,18,19,27,28,29,31,32  
 Seq#:4; Xaa Pos. 33,34,35,36,37,38,39  
 Seq#:5; Xaa Pos. 1,2,3,5,6,7,9,10,13,14,16,28,30,31,32,33,34  
 Seq#:6; Xaa Pos. 5,14,26,31  
 Seq#:7; Xaa Pos. 1,3,7,16,31,32,34  
 Seq#:8; Xaa Pos. 1,2,4,10,19,31,34,36,37,39  
 Seq#:9; Xaa Pos. 9,13,17  
 Seq#:10; Xaa Pos. 2,3,10,12,18,26,32,33  
 Seq#:11; Xaa Pos. 3,5,16,18,22,25,31  
 Seq#:12; Xaa Pos. 1,5,8,11  
 Seq#:13; Xaa Pos. 5,8,10,15,16,23  
 Seq#:14; Xaa Pos. 4,9,12,13,17  
 Seq#:15; Xaa Pos. 8,15,16  
 Seq#:24; N Pos. 17  
 Seq#:36; Xaa Pos. 3,7,8,14,18  
 Seq#:37; Xaa Pos. 1  
 Seq#:39; N Pos. 12  
 Seq#:41; N Pos. 6,9,12,15,18,21,24  
 Seq#:42; Xaa Pos. 14,26,31  
 Seq#:43; Xaa Pos. 9,13  
 Seq#:44; Xaa Pos. 2  
 Seq#:45; Xaa Pos. 1,6,7,8,12,17,18,23

## VERIFICATION SUMMARY

DATE: 02/13/2004

PATENT APPLICATION: US/10/647,519

TIME: 12:04:45

Input Set : A:\2314-269seq.txt

Output Set: N:\CRF4\02132004\J647519.raw

L:21 M:270 C: Current Application Number differs, Replaced Application Number

L:22 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:72 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0

M:341 Repeated in SeqNo=1

L:124 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0

M:341 Repeated in SeqNo=2

L:166 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0

M:341 Repeated in SeqNo=3

L:208 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0

M:341 Repeated in SeqNo=4

L:250 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0

M:341 Repeated in SeqNo=5

L:274 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0

M:341 Repeated in SeqNo=6

L:298 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0

M:341 Repeated in SeqNo=7

L:322 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0

M:341 Repeated in SeqNo=8

L:344 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0

M:341 Repeated in SeqNo=9

L:365 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0

M:341 Repeated in SeqNo=10

L:389 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0

M:341 Repeated in SeqNo=11

L:409 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0

L:433 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:0

M:341 Repeated in SeqNo=13

L:453 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:0

M:341 Repeated in SeqNo=14

L:473 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0

L:734 M:258 W: Mandatory Feature missing, &lt;223&gt; Tag not found for SEQ ID#:24

L:734 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0

L:1091 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 after pos.:0

M:341 Repeated in SeqNo=36

L:1113 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:0

L:1140 M:258 W: Mandatory Feature missing, &lt;221&gt; Tag not found for SEQ ID#:39

L:1140 M:258 W: Mandatory Feature missing, &lt;222&gt; Tag not found for SEQ ID#:39

L:1140 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39 after pos.:0

L:1168 M:258 W: Mandatory Feature missing, &lt;221&gt; Tag not found for SEQ ID#:41

L:1168 M:258 W: Mandatory Feature missing, &lt;222&gt; Tag not found for SEQ ID#:41

L:1168 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41 after pos.:0

L:1184 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:42 after pos.:0

M:341 Repeated in SeqNo=42

L:1206 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:43 after pos.:0

L:1224 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:44 after pos.:0

L:1241 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45 after pos.:0

M:341 Repeated in SeqNo=45